

CRF Errors Corrected by the STIC Systems Branch

OIR
11/5/98

Serial Number: 09/177,843

CRF Processing Date: 11/5/98
Edited by: AK
Verified by: _____ (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☒ Other: Seq 2 - removed the answer and

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/177,843DATE: 11/09/98
TIME: 16:52:25

INPUT SET: S29699.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

 ENTERED

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: THE TRUSTEES OF COLUMBIA UNIVERSITY IN THE
CITY OF NEW YORK(ii) TITLE OF INVENTION: A METHOD OF PREVENTING AND TREATING
BACTERIAL INFECTION OF SUTURES AND
PROSTHETIC DEVICES, AND PROMOTING
INGRESS OF LEUKOCYTES INTO TUMOR
FOCI

(iii) NUMBER OF SEQUENCES: 2

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Cooper & Dunham LLP
(B) STREET: 1185 Avenue of the Americas
(C) CITY: New York
(D) STATE: New York
(E) COUNTRY: U.S.A.
(F) ZIP: 10036

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: Not Yet Known
(B) FILING DATE: April 22, 1997
(C) CLASSIFICATION:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: White, John P
(B) REGISTRATION NUMBER: 28,678
(C) REFERENCE/DOCKET NUMBER: 48940-A-PCT/JPW/JKM

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 212-278-0400
(B) TELEFAX: 212-391-0525

(2) INFORMATION FOR SEQ ID NO:1:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/177,843DATE: 11/09/98
TIME: 16:52:27

INPUT SET: S29699.raw

47 (i) SEQUENCE CHARACTERISTICS:
48 (A) LENGTH: 6 amino acids
49 (B) TYPE: amino acid
50 (C) STRANDEDNESS: single
51 (D) TOPOLOGY: linear
52
53 (ii) MOLECULE TYPE: peptide
54
55 (iii) HYPOTHETICAL: NO
56
57 (iv) ANTI-SENSE: NO
58
59
60
61
62
63 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
64
65 Gly Arg Gly Asp Ser Pro
66 1 5
67
68
69 (2) INFORMATION FOR SEQ ID NO:2:
70
71 (i) SEQUENCE CHARACTERISTICS:
72 (A) LENGTH: 6 amino acids
73 (B) TYPE: amino acid
74 (C) STRANDEDNESS: single
75 (D) TOPOLOGY: linear
76
77 (ii) MOLECULE TYPE: peptide
78
79 (iii) HYPOTHETICAL: NO
80
81 (iv) ANTI-SENSE: NO
82
83
84
85
86 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
87
88 Gly Arg Gly Glu Ser Pro
89 1 5
90

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/177,843

DATE: 11/09/98
TIME: 16:52:28

INPUT SET: S29699.raw

Line	Error	Original Text
31	Wrong application Serial Number	(A) APPLICATION NUMBER: Not Yet Known

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/177,843DATE: 11/05/98
TIME: 18:55:20

INPUT SET: S29699.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

(1) General Information:

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INGRESS OF LEUKOCYTES INTO TUMOR
FOCI

(iii) NUMBER OF SEQUENCES: 2

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(B) TELEFAX: 212-391-0525

(2) INFORMATION FOR SEQ ID NO:1:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/177,843DATE: 11/05/98
TIME: 18:55:22

INPUT SET: S29699.raw

47 (i) SEQUENCE CHARACTERISTICS:
48 (A) LENGTH: 6 amino acids
49 (B) TYPE: amino acid
50 (C) STRANDEDNESS: single
51 (D) TOPOLOGY: linear
52
53 (ii) MOLECULE TYPE: peptide
54
55 (iii) HYPOTHETICAL: NO
56
57 (iv) ANTI-SENSE: NO
58
59
60
61
62
63 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
64
65 Gly Arg Gly Asp Ser Pro
66 1 5
67
68
69 (2) INFORMATION FOR SEQ ID NO:2:
70
71 (i) SEQUENCE CHARACTERISTICS:
72 (A) LENGTH: 6 amino acids
73 (B) TYPE: amino acid
74 (C) STRANDEDNESS: single
75 (D) TOPOLOGY: linear
76
77 (ii) MOLECULE TYPE: peptide
78
79 (iii) HYPOTHETICAL: NO
80
81 (iv) ANTI-SENSE: NO
82
83
84
85
86 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
87
88 Gly Arg Gly Glu Ser Pro
89 1 5

number the amino acids

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/177,843

DATE: 11/05/98
TIME: 18:55:24

INPUT SET: S29699.raw

Line	Error	Original Text
31	Wrong application Serial Number	(A) APPLICATION NUMBER: Not Yet Known